1600

```
        RAW SEQUENCE LISTING              DATE: 09/23/2003
        PATENT APPLICATION: US/09/618,361B   TIME: 18:27:59

        Input Set : A:\PTO.KD.txt
        Output Set: N:\CRF4\09232003\I618361B.raw

   2 <110> APPLICANT: Balasubramanium, Ambikaipakan
   3        Chance, William T.
   5        University of Cincinnati
   7 <120> TITLE OF INVENTION: Compounds for Control of Appetite, Blood Pressure,
   8        Cardiovascular Response, Libido and Circadian Rhythm
  10 <130> FILE REFERENCE: UOC-136R
  12 <140> CURRENT APPLICATION NUMBER: US/09/618,361B
  13 <141> CURRENT FILING DATE: 2000-07-18
  15 <150> PRIOR APPLICATION NUMBER: US/09/449,914
  16 <151> PRIOR FILING DATE: 1999-12-02
  18 <160> NUMBER OF SEQ ID NOS: 7
  20 <170> SOFTWARE: PatentIn Version 3.1
```

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

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      36 <210> SEQ ID NO: 2
      37 <211> LENGTH: 4
      38 <212> TYPE: PRT
      39 <213> ORGANISM: Artificial Sequence
      41 <220> FEATURE:
W-->  42 <221> NAME/KEY: MOD RES
      43 <222> LOCATION: (1)...(4)
      44 <223> OTHER INFORMATION: The compound is a dimer of the sequence listing
      45        and the dimer is formed by a disulfide bond. Artificial
      46        sequence is completely synthesized.
E-->  48 <400> SEQUENCE: 2
      49 Cys Trp Arg Tyr
      50  1
      97 <210> SEQ ID NO: 6
      98 <211> LENGTH: 6
      99 <212> TYPE: PRT
     100 <213> ORGANISM: Homo sapiens
     102 <400> SEQUENCE: 6
     103 Trp Arg Tyr Trp Arg Tyr
E--> 104  1                       5 <- insert
     109 <210> SEQ ID NO: 7
     110 <211> LENGTH: 6
     111 <212> TYPE: PRT
     112 <213> ORGANISM: Homo sapiens
E--> 114 <400> SEQUENCE: 7
     115 Trp Arg Tyr Trp Arg Tyr
     116  1               5
```

VERIFICATION SUMMARY   DATE: 09/23/2003
PATENT APPLICATION: **US/09/618,361B**   TIME: 18:28:00

Input Set : **A:\PTO.KD.txt**
Output Set: **N:\CRF4\09232003\I618361B.raw**

```
L:42 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:48 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:4
L:61 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:67 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:87 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:93 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:104 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:114 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:7 differs:6
```